



SEQUENCE LISTING

<110> HERING, THOMAS M.
JOHNSTONE, BRIAN

<120> PROBES FOR CHONDROGENESIS

<130> 27708/04065

<140> 10/623,914
<141> 2003-07-21

<150> 60/211,384
<151> 2000-06-14

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<170> PatentIn Ver. 3.3

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<212> DNA

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 Lys Lys Val Met Leu Glu Asn Tyr Arg Asn Leu Val Ser Val Gly Leu
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 Ile Thr His Ile Asp Thr Leu Ile Glu Lys Arg Asp His Ser Asn Lys
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 Ser Gly Thr Val Phe His Leu Asn Thr Leu Ser Tyr Ile Lys Gln Ile
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 Phe Pro Met Glu Glu Arg Ile Phe Asn Phe His Thr Asp Lys Ser
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 Val His Thr Gly Glu Lys Pro Tyr Gln Cys Lys Gln Cys Asn Lys Ala
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 Glu Lys Pro Tyr Glu Cys Ile Glu Cys Gly Lys Ala Phe Ser Asp Cys
 340 345 350
 Ser Ser Leu Ala His His Arg Arg Ile His Thr Gly Lys Arg Pro Tyr
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 Glu Cys Ile Asp Cys Gly Lys Ala Phe Arg Gln Asn Ala Ser Leu Ile
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 Arg His Arg Arg Tyr Tyr His Thr Gly Glu Lys Pro Phe Asp Cys Ile
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 Arg Ile His Thr Gly Glu Arg Pro Tyr Lys Cys Asn Val Cys Gly Lys
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 Arg Gly Ser Leu Thr Leu His Gln Arg Val His Thr Gly Glu Lys Pro
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 Gly Glu Lys Pro Tyr Glu Cys Ile Glu Cys Gly Lys Ala Phe Ser Asp
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 Gly Ser Tyr Leu Val Gln His Pro Arg Leu His Ser Gly Lys Arg Pro
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 Tyr Glu Cys Leu Glu Cys Gly Lys Ala Phe Arg Gln Arg Ala Ser Leu
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 Val Cys Gly Lys Ala Phe Ser His Arg Lys Ser Leu Thr Leu His Gln
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 Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Glu Cys Ser Lys
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Ser Lys Thr Tyr Glu Thr Lys Ile Phe Ser Glu Asn Asp Ile Phe 50 55 60
Glu Ile Asn Phe Ser Gln Trp Glu Met Lys Asp Lys Ser Lys Thr Leu 65 70 75 80
Gly Leu Glu Ala Ser Ile Phe Arg Asn Asn Trp Lys Cys Lys Ser Ile 85 90 95
Phe Glu Gly Leu Lys Gly His Gln Glu Gly Tyr Phe Ser Gln Met Ile 100 105 110
Ile Ser Tyr Glu Lys Ile Pro Ser Tyr Arg Lys Ser Lys Ser Leu Thr 115 120 125
Pro His Gln Arg Ile His Asn Thr Glu Lys Ser Tyr Val Cys Lys Glu 130 135 140

Cys Gly Lys Ala Cys Ser His Gly Ser Lys Leu Val Gln His Glu Arg
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 Thr His Thr Ala Glu Lys His Phe Glu Cys Lys Glu Cys Gly Lys Asn
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 Tyr Leu Ser Ala Tyr Gln Leu Asn Val His Gln Arg Phe His Thr Gly
 180 185 190
 Glu Lys Pro Tyr Glu Cys Lys Glu Cys Gly Lys Thr Phe Ser Trp Gly
 195 200 205
 Ser Ser Leu Val Lys His Glu Arg Ile His Thr Gly Glu Lys Pro Tyr
 210 215 220
 Glu Cys Lys Glu Cys Gly Lys Ala Phe Ser Arg Gly Tyr His Leu Thr
 225 230 235 240
 Gln His Gln Lys Ile His Ile Gly Val Lys Ser Tyr Lys Cys Lys Glu
 245 250 255
 Cys Gly Lys Ala Phe Phe Trp Gly Ser Ser Leu Ala Lys His Glu Ile
 260 265 270
 Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Lys Glu Cys Gly Lys Ala
 275 280 285
 Phe Ser Arg Gly Tyr Gln Leu Thr Gln His Gln Lys Ile His Thr Gly
 290 295 300
 Lys Lys Pro Tyr Glu Cys Lys Ile Cys Gly Lys Ala Phe Cys Trp Gly
 305 310 315 320
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 325 330 335
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 Gln His Glu Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Glu
 355 360 365
 Cys Gly Lys Ala Phe Ser Arg Gly Tyr His Leu Ser Gln His Gln Lys
 370 375 380
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 385 390 395 400
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 405 410 415
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 420 425 430
 Tyr Gln Leu Thr Arg His Gln Val Phe His Thr Gly Glu Lys Pro Tyr
 435 440 445

Glu Cys Lys Glu Cys Gly Lys Ala Phe Asn Cys Gly Ser Ser Leu Val
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 Gln His Glu Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Glu
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